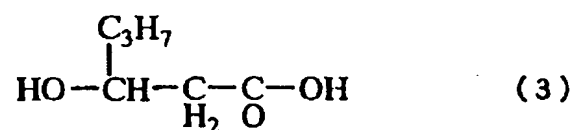
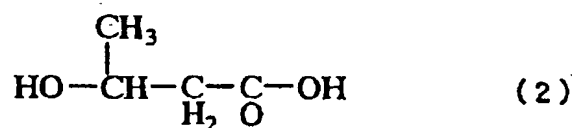


in the formula, R represents an alkyl group.

3. The transformant according to Claim 1

wherein the polyester is copolyester P(3HB-co-3HH) resulting from the copolymerization of 3-hydroxybutyric acid of the following formula (2) and 3-hydroxyhexanoic acid of the following formula (3);



4. The transformant according to claim 1

wherein the yeast belongs to any of the genera

Aciculoconidium, Ambrosiozyma, Arthroascus, Arxiozyma, Ashbya, Babjevia, Bensingtonia, Botryoascus, Botryozyma, Brettanomyces, Bullera, Bulleromyces, Candida, Citeromyces, Clavispora, Cryptococcus, Cystofilobasidium, Debaryomyces, Dekkara, Dipodascopsis, Dipodascus, Eeniella, Endomycopsella, Eremascus, Eremothecium, Erythrobasidium, Fellomyces, Filobasidium, Galactomyces, Geotrichum, Guilliermondella, Hanseniaspora, Hansenula, Hasegawaea, Holtermannia, Hormoascus, Hyphopichia, Issatchenkia, Kloeckera, Kloeckeraspora, Kluyveromyces, Kondoa, Kuraishia, Kurtzmanomyces, Leucosporidium, Lipomyces, Lodderomyces, Malassezia, Metschnikowia, Mrakia, Myxozyma, Nadsonia, Nakazawaea, Nematospora, Ogataea, Oosporidium, Pachysolen, Phachytichospora, Phaffia, Pichia, Rhodospiridium, Rhodotorula, Saccharomyces, Saccharomycodes, Saccharomycopsis, Saitoella, Sakaguchia, Saturnospora, Schizoblastosporion, Schizosaccharomyces, Schwanniomyces, Sporidiobolus, Sporobolomyces, Sporopachydermia, Stephanoascus, Sterigmatomyces, Sterigmatosporidium, Symbiotaphrina, Sympodiomyces, Sympodiomycopsis, Torulaspora, Trichosporiella, Trichosporon, Trigonopsis, Tsuchiyaea, Udeniomyces, Waltomyces, Wickerhamia, Wickerhamiella, Williopsis, Yamadazyma, Yarrowia, Zygoascus, Zygosaccharomyces, Zygowilliopsis and Zygozyma.

5. The transformant according to claim 1
wherein the yeast is Yarrowia lipolytica.
6. The transformant according to claim 1
wherein the yeast is Candida maltosa.
7. The transformant according to claim 1
wherein a polyester synthesis-associated enzyme gene expression cassette
comprises a promoter and a terminator,
said promoter and said terminator functioning in a yeast.
8. The transformant according to Claim 7
wherein the promoter and terminator are derived from Yarrowia lipolytica.
9. The transformant according to Claim 7
wherein the promoter is derived from Yarrowia lipolytica ALK3.
10. The transformant according to Claim 7
wherein the terminator is derived from Yarrowia lipolytica XPR2.
11. The transformant according to Claim 7
wherein the promoter and terminator are derived from Candida maltosa.
12. The transformant according to Claim 7
wherein the promoter is derived from Candida maltosa ALK1.
13. The transformant according to Claim 7
wherein the terminator is derived from Candida maltosa ALK1.
14. The transformant according to Claim 1
wherein the polyester synthesis-associated enzyme gene is derived from
Aeromonas caviae.
15. The transformant according to Claim 1

wherein the polyester synthesis-associated enzyme gene is a PHA synthase gene derived from Aeromonas caviae or a PHA synthase gene and (R)-specific enoyl-CoA hydratase gene.

16. The transformant according to Claim 15
wherein said PHA synthase gene has the sequence represented by SEQ ID N0:3
and the (R)-specific enoyl-CoA hydratase gene has the sequence represented by SEQ ID N0:4.
17. A method of producing a polyester using the transformant according to Claim 1
which comprises growing said transformant and harvesting a polyester from the resulting culture.
18. A polyester synthesis-associated enzyme gene which is modified from at least one gene code CTG to TTA, TTG, CTT, CTC or CTA.
19. The polyester synthesis-associated enzyme gene according to Claim 18
which codes for an enzyme derived from a bacterium.
20. The polyester synthesis-associated enzyme gene according to Claim 19
wherein said bacterium is Aeromonas caviae.
21. The polyester synthesis-associated enzyme gene according to Claim 20
wherein the enzyme gene derived from Aeromonas caviae is a PHA synthase gene or a (R) -specific enoyl-CoA hydratase gene.
22. The polyester synthesis-associated enzyme gene according to Claim 21
wherein said PHA synthase gene has the sequence represented by SEQ ID N0:3.
23. The polyester synthesis-associated enzyme gene according to Claim 21
wherein said (R)-specific enoyl-CoA hydratase gene has the sequence represented by SEQ ID N0:4.